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3: /cgn2_6/ptodata/2,
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1 MTSPSSSPVFRLET
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July 18, 2001, 15:54:31 ; Search time 13.86 Seconds (without alignments) 1110.426 Million cell updates/sec
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Copyright (c) 1993 - 2000 Compugen Ltd
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US-08-623-679-9

US-08-623-679-9

US-08-933-774-9

US-09-031-485-23

US-09-065-474-23

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US-09-065-474-23

US-09-08-477-429A-28

US-09-08-474-28

US-08-847-429A-28

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US-09-108-93-33
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US-08-185-432-19
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US-09-197-636-2
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Result

No

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	Sequence 4,	Sequence 4,	Sequence 2,	Sequence 2,	Sequence 2,	Sequence 4,	Sequence 2,	Sequence 4,	Sequence 2,	Sequence 4,	Sequence 2,	Sequence 2,	Sequence 38	Sequence 38	Sequence 38	Sequence 2,	Sequence 20	Sequence 20
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ALIGNMENTS

; MOLECULE TYPE: US-09-197-636-2 Sequence 2, Application US/09197636 Patent No. 6239267 APPLICATION NUMBER: US/09/197,636
FILING DATE: 23-NOV-1998
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: UK 9805137.8
FILING DATE: 12-MAR-1998
APPLICATION NUMBER: UK 9815791.0
FILING DATE: 21-JUL-1998
APPLICATION NUMBER: UK 9819278.4
FILING DATE: 03-SEP-1998
APPLICATION NUMBER: UK 9819278.4
FILING DATE: 03-SEP-1998
ATTORNEY/AGENT INFORMATION:
NAME: PRESCIA, Paul F GENERAL INFORMATION:

APPLICANT: DUCKWORTH, DAVI
APPLICANT: HAVES, PHILIP
APPLICANT: MEADOWS, HELEN INFORMATION FOR SEQ ID NO: COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA: TITLE OF INVENTION: NOVEL COMPOUNDS NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS: SEQUENCE CHARACTERISTICS: LENGTH: 839 amino acids REGISTRATION NUMBER: 23,031
REFERENCE/DOCKET NUMBER: GP
TELECOMMUNICATION INFORMATION:
TELEPHONE: 601.407-0700 APPLICANT: STREET: P.O. Box 980
CITY: Valley Forge
STATE: PA TELEPHONE: 601-40/-0/ TELEFAX: 610-407-0701 TELEX: 846169 TOPOLOGY: TYPE: amino acid STRANDEDNESS: sin OPERATING SYSTEM: COUNTRY: 19482-0980 SD DAVIS, JOHN DUCKWORTH, DAVID single US/09/197,636

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US-09-197-636-8
; Sequence 8, Application US/09197636
; Patent No. 6239267
; GENERAL INFORMATION:
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Best Local Similarity
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APPLICANT:
APPLICANT:
                                                                                             TITLE OF INVENTION: NOVEL COMPOUNDS
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS.
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                                CITY: Valley Forge
STATE: PA
                  COUNTRY:
                                                                    ADDRESSEE: Ratner & Prestia STREET: P.O. Box 980
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MEADOWS, HELEN
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48.5%; Pred. No. 9.10
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches 344;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING CLASSIFICATION:
CLASSIFICATION DATA:
PRIOR APPLICATION NUMBER: UK
APPLICATION NUMBER: 12-MAR-190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 12-MAR-1998
APPLICATION NUMBER: UK 9815791.0
FILING DATE: 21-UL-1998
APPLICATION NUMBER: UK 9819278.4
FILING DATE: 03-SEP-1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: 601-407-0700
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APPLICATION NUMBER: US/09/197,636
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MEDIUM TYPE: Diskette
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TELEFAX: 610
TOTEX: 846169
                                  548
590
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REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OPERATING SYSTEM:
SOFTWARE: FastSEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                               GFSTAVVTLIEDGKNDSLPSESTSHRWRGPACRPPDSS
                                GFAVALVSLSQEA----
                                                               YFSHLKEYVASMVFSLALGWINMLYYTRGFQQMGIYAVMIEKMILRDLCRFMFVYVVFLF
                                                                                  CFLAIEWYLPLLVSALVLGWLNLLYYTRGFQHTGIYSVMIQKVILRDLLRFLLIYLVFLF 547
                                                                                                                                                                     GNSMLLTGHILILLGGIYLLVGQLWYFWRRHVFIWISFIDSYFEILFLFQALLTVVSQVL 487
                                                                                                                                                                                                                                                                          GDYFRVTGEILSVLGGVYFFFRGIQYFLQRRPSMKTLFVDSYSEMLFFLQSLFMLATVVL
                                                                                                                                                                                                                                          MVVLEPLNKLLQAKWDLLIPK-FFLNFLCNLIYMFIFTAVAYHQPTLKKQAAPHLKAE-V 427
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TIPLLLEIARQTDSLKELVNASYTDSYYKGQTALHIAIERRNMALVTLLVENGADVQAAA 232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            YDRRSIFEAVAQNNCQDLESLLLFLQKSKKHLTDNEFKDPETGKTCLLKAMLNLHDGQNT 172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     amino acid
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FastSEQ for Windows
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                                                                                       Best Local Similarity
                                                                                                     Query Match
                                                                                                                                                                                                                                                                         TELEX: 846169
INFORMATION FOR SEQ ID NO: 4:
                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: UK 980
FILING DATE: 12-MAR-1998
APPLICATION NUMBER: UK 981
FILING DATE: 21-JUL-1998
APPLICATION NUMBER: UK 981
FILING DATE: 02-CFD-1000
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                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 601-407-0700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 03-SEP-1998
ATTORNEY/AGENT INFORMATION:
NAME: Prestia, Paul F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: NOVEL COMPOUNDS NUMBER OF SEQUENCES: 8
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     113
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                 74 FDRDRLFNAVSRGVPEDLAGLPEYLSKTSKYLTDSEYTEGSTGKTCLMKAVLNLKDGVNA 133
                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: si
TOPOLOGY: linear
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TELEPHONE: 610-407-0701
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STREET: P.O. Box 980
CITY: Valley Forge
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OPERATING SYSTEM:
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SOFTWARE: FastSEQ for Windows Version
                                                                                                                                                                                                                                           LENGTH:
YDRRSIFEAVAQNNCQDLESLLLFLQKSKKHXTDNEFKDPETGKTCLLKAMLNLHDGQNT 172
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DAVIS, JOHN
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                                                                     Conservative 121; Mismatches 201;
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                                                                                                                                                                                                       single
                                                                                   40.9%; Score 1638.5; DB 4 48.3%; Pred. No. 4.2e-146;
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                                                                                                     DB 4;
                                                                   Indels
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; OTHER INFORMATION: 929491 US-09-172-977-4
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                                                                                   NUMBER OF SEQ ID NOS: 4
SOFTWARE: PERL Program
SEQ ID NO 4
LENGTH: 1839
                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                              Sequence 4, Application US/09172977 Patent No. 5989863
                                                                                                                                                          APPLICANT: Tang, Y. Tom
APPLICANT: Guegler, Karl J.
APPLICANT: Corley, Neil C.
APPLICANT: Yue, Henry
TITLE OF INVENTION: HUMAN ANKYRIN FAMILY PROTEIN
FILE REFERENCE: PF-0615 US
CURRENT APPLICATION NUMBER: US/09/172,977
CURRENT FILING DATE: 1998-10-14
                              TYPE: PRT
ORGANISM: Homo sapiens
FEATURE: -
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Query Match
Best Local Similarity
Matches 88; Conserv

Conservative

46;

Score 151; DB 2; Pred. No. 6.9e-05; 6; Mismatches 96;

Length 1839; Indels

92;

Gaps

19;

3.8%;

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Gaps

19;

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US-09-172-977-3
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US-09-082-059-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; ORGANISM: Homo sapiens US-09-082-059-2
                                                                      GENERAL INFORMATION:
                                                                                         Sequence 3, Application US/09172977 Patent No. 5989863
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 2
LENGTH: 1088
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 2, Application US/09082059A Patent No. 6225086 GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/082
CURRENT FILING DATE: 1998-05-21
EARLIER APPLICATION NUMBER: 60/047356
EARLIER FILING DATE: 1997-05-21
NUMBER OF SEQ. ID NOS: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Morrow, Jon S.

APPLICANT: Devarajan, Prasad
TITLE OF INVENTION: No. 6225086el Ankyrin Proteins and a Method for Their Identificat
FILE REFERENCE: 44574-5002-US.
APPLICANT:
                        APPLICANT:
                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
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                                                                                                                                                                                                                                                          282 PTVQLEDIRNLQDLTPLKLAAKEGKIEIFRHILQREFS 319
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                                                                                                                                                                                                                                  181 ITTK-----KGFTPLHVAAKYGKLEVANLLLQKSAS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     KKQAAPHLKAEVGNSMLLTGHI 437
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             --MAHPDAATTNG-YTPLHISAREGQVDVASV-----LLEAGAAHSLATKKG----FT 565
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SAENIALVTSMYDGLLQAGARLCPTVQLEDIRNLQDLTPLKLAAKEGKIEIFRHILQREF 318
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                                                                                                                                                                                                                                                                                                                                                                          WDVVSYLLENPHQPASLQATDSQGNTVLHALVMISDNSAENIALVTSMYDGLLQAGARLC 281
                                                                                                                                                                                                                                                                                                                                                                                                                        RGETALHMAARSGQAEVVRYLVQDGAQVEAKA-----KDDQT------PLHISARLGK 132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RGHSALHIAIEKRSLQCVKLLVENGANVHARACGRFFQKGQGTCFYFGELPLSLAACTKQ 221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PLNKLLQAKWDLL-IPKFFL----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RAGQVEVVRC----LLRNGA-----LVDARAREEQTPLHIASRLGKTEIVQLLLQH-- 519
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             QKGQGTCFYFGELPLSLAACTKQWDVVSYLLENPHQPASLQATDSQGNTVLHALVMISDN 258
                                                                                                                                                                                                                                                                                                                              ADIVQQLLQ---QGASPNAATTSGYTPLH---LSAREGHEDVAAF---
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PLH--VAAKYGSLDVAKLLLQRRAAADSAGKNGLTPL-----HVAAHYDNQKVALLLL 616
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -----ITESGLTPIHVAAFMGHLNIVLLLLQN---GASPDVTNIRGETALHMAA-----
Corley, Neil C
                     Tang, Y. Tom
Guegler, Kar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                        Karl J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.7%; Score 150; DB 4; Length 1088; 32.3%; Pred. No. 3.8e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US/09/082,059A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               --NFLCNLIYMFIFTAVAYHQPT----L 415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    51;
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                                                                                                                                                                                                                                                                                                                              LLDHGASLS 180
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-112-096-15; Sequence 15, Application US/09112096; Patent No. 6194152
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         Query Match
Best Local Similarity
Matches 151; Conserv
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TITLE OF INVENTION: HUMAN ANKYRIN FAMILY PROTEIN
FILE REFERENCE: PF-0615 US
CURRENT APPLICATION NUMBER: US/09/172,977
CURRENT FILING DATE: 1998-10-14
NUMBER OF SEQ ID NOS: 4
SOFTWARE: PERL Program
SEQ ID NO 3
                                                                                                                                                                                                       SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 15 .
LENGTH: 1095
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                                                                                                                                                                             EARLIER APPLICATION NUMBER: 60/056,110 EARLIER FILING DATE: 1997-08-20 NUMBER OF SEQ ID NOS: 29
                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Prostate Tumor Polynucleotide TITLE OF INVENTION: Antigen Compositions FILE REFERENCE: 7636-0015.30
CURRENT APPLICATION NUMBER: US/09/112.096
CURRENT FILING DATE: 1998-07-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Reiner Laus
APPLICANT: Michael H. Shapero
APPLICANT: Larisa Tsavaler
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 843
TYPE: PRT
                                                                                                                                                           ORGANISM: Homo sapiens
                                                                                                                                                                                      TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Rattus norvegicus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  416 KKQAAPHLKAEVGNSMLLTGHI 437
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  531 PLH--VAAKYGSLDVAKLLLQRRAAADSAGKNGLTPL-----HVAAHYDNQKVALLLL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      139 LQIDRDSGNPQPLVNAQÇTDDYYRGHSALHIAIEKRSLQCVKLLVENGANVHARACGRFF 198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   392 -----ITESGLTPIPVAAFMGHLNIVLLLLQN---GASPDVTNIRGETALHMAA----- 437
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EKGASPHATAKNGYTPL---HI 600
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RAGEVEVVRC----LLRNGA-----LVDARAREEQTPLHIASRLGKTEIVQLLLQH--
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                     Conservative
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                                         3.6%;
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27.3%; Pred. No. 3.9e-05;
27.3%; Mismatches 97;
                     126;
                                         Score 146; DB 4;
Pred. No. 9.1e-05;
                     Mismatches
                                                            Length 1095;
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RESULT 8
US-09-065-474-139
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                                                                                                                                              Sequence 139, Application US/09065474 Patent No. 6063599
                                                                                                                            GENERAL INFORMATION:
         APPLICANT: Tang, Liang
APPLICANT: Blehm, E. Scot
TITLE OF INVENTION: DIROFILARIA AND BRUGIA ANKYRIN
TITLE OF INVENTION: PROTBINS, NUCLEIC ACID MOLECULE
TITLE OF INVENTION: USES THEREOF
NUMBER OF SEQUENCES: 171
                                                                                                                                                                                                                                                                1042
CORRESPONDENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          530
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                                                                                                                                                                                                                                                                                                  NWASWE 715
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IEKRSLQCVKLLVENGANVHARACGRFFQKGQGTCFYFGELPLSLAACTKQWDVVSYLLE 230
                                                                                                                                                                                                                                                              ETLAWE 1047
                                                                                                                                                                                                                                                                                                                                     YCSRLNIPFPFIVFAYFYMVVKKCFKCCCKEKNMESSVC--
                                                                                                                                                                                                                                                                                                                                                                           FGQVPSDVDGTTYDFAHCTFTGNESKPLCVELDEHN---
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           -PEAPTGPNAT-----ESVOPMEGQEDEGNGAQYRGILEASLELFKFTIGMGEL 610
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LGPKIIMLQRMLI-DVFFFLFLFAVWMVAFGVARQGILRQNEQRWRWIFRSVIYEPYLAM 902
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -----VMIQKVILRDLLRFLLIYLVFLFGFAVA---LVSLSQEAWR-------
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       YFTDLWNVMDTLGLFYFIAGIVERLHSSNKSSLYSGRVIFCLDYIIFTLRLIHIFTVSRN 843
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -----VLGWLNLLYYTRGF------QHTGIYS---------
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    -WRRHVFIWISFIDSYFEILFL-FQA------LLTVVSQVLCFLAIEWYLPLLVSAL- 503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SKQWYGEISRDTKNWKIILCLFIIPLVGCGFVSFRKKPVDKHKKLLWYYVAFFTSPFVVF 729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 KKQAAPHLKAEVGN-SMLLTGHILILLGGIYLLVGQ-----LWYF-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CEA-----GSNCLELAVEATDQHFTAQPGVQNFL 669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CKSPHRHRMVVLEPLNKLLQAKWDLLIPKFFLNFLCNLIYMFI-----FTAVAYHQPTL 415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TLAALGASKLLKTLAKVKNDINAAGESEELANEYETRAVELFTECYSSDEDLAEQLLVYS 638
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   VRVSL-----SCEENSVLEIIAFH 361
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DRNGRDEMDIELHDVSP-----ITRHPLQALFIWAILQNKKELSKVIWEQTRGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IRN------LQDLTPLKLAAKEGKIEIFRHILQREF-----SGLSHLSRKFTEWCYGP 335
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NPHQPASLQATDSQGNTVLHALVMISDNSAENIALVTSMYD--GLLQAGARLCPTVQLED 288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NACILPLLQI-----DRDSGNPQ-----PLVNAQCTDDYYRGHSA-----LHIA 170
                                                                                                                                                                                                                                                                                                                                                                                                              -----EWITIPLVCIYMLSTNILLVNLLVAMFGYTVGTVQENNDQVWKFQRYFLVQE 995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SW--NVVFYIAFLLLFAYVLLMDFHSVPHPPELVLYSLVFVLFCDEVRQWY----VNGVN 783
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 N------HFSTLVYRNLQIAKNS-YNDALLTFVWKLVANFRRGFR-----KE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SRGVPEDLAG-LPEYLSKTSKYLTDS-----EYTEGSTGKTCLMKAVLNLKDG-----V
ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                           -----CRKKQRAGVMLTVGTKPDGSPDERWCFRVEEV 709
                                                   MOLECULES,
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US-09-031-485-33; Sequence 33, Application US/09031485; Patent No. 5824306; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Verser, Carol Talkington
REGISTRATION NUMBER: 37,459
REFERENCE/DOCKET NUMBER: HW-5-C1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 970/493-7272
TELEFAX: 970/484-9505
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 139:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: WordPerfect for Windows,
CURRENT APPLICATION DATA:
                                                                                                                                                   APPLICANT: Tang, Liang
APPLICANT: Blehm, E. Scot
APPLICANT: Blehm, E. Scot
TITLE OF INVENTION: DIROFILARIA AND BRUGIA ANKYRIN
TITLE OF INVENTION: PROTEINS, NUCLEIC ACID MOLECULES, AND
TITLE OF INVENTION: USES THEREOF
NUMBER OF SEQUENCES: 85
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       227 YLLENPHQPASLQATDSQGNTVLHALVMISDNSAENIALVTSMYDGLLQAGARLCPTVQL 286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  167 LHIAIEKRSLQCVKLLVENGANVHARACGRFFQKGQGTCFYFGELPLSLAACTKQWDVVS 226
                                                                          STREET: 1825 Sharp CITY: Fort Collins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: FILING DATE: 24-APPLICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 1825 Sharp
CITY: Fort Collins
                                                        STATE:
                                                                                                               ADDRESSEE: Heska Corporation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Carol Talkington Verser, ADDRESSEE: Heska Corporation STREET: 1825 Sharp Point Drive
                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TEWCYGPVRVSLYDLASVDSCEE-----NSVLEIIAFHCKSPHRHRWVVLEPLN 377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LHVAAHYNNDKVALLLLENGASAHAAAKN------GYTPLHIAAKKNQMDIAS 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -----NGLTPMHLCAQEDRVSVAEELVKENAAIDPKTKAGYTPLHVACHFGQINMVRFL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EDIRNLQDLTPLKLAAKEGKIEIFRHILQREFS-------GLSHLSRKF 328
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                                                        Colorado
                                                                                             1825 Sharp Point Drive
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MEDIUM TYPE:

US/08/847,429A

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; TOPOLOGY: linear; MOLECULE TYPE: protein US-09-031-485-33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 33, Application US/08847429A Patent No. 5827692
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 33:
                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: Windows 95
                                                                                                                                                                                                                                                                                                                             APPLICANT: Blehm, E. Scot
TITLE OF INVENTION: DIROFILARIA AND BRUGIA ANKYRIN
TITLE OF INVENTION: PROFEINS, NUCLEIC ACID MOLECULE
TITLE OF INVENTION: USES THEREOF
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 1745 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
FILING DATE: 24-APR-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               730 IE--HG-ARVSVITRASYTPLHQAAQQGHNSVVRYLLEHGASPNVHTSTGQTPLS 781
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        329 TEWCYGPVRVSLYDLASVDSCEE-----NSVLEIIAFHCKSPHRHRMVVLEPLN 377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  227 YLLENPHQPASLQATDSQGNTVLHALVMISDNSAENIALVTSMYDGLLQAGARLCPTVQL 286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                582 LHVAAHYNNDKVALLLLENGASAHAAAKN------GYTPLHIAAKKNQMDIAS 628
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          167 LHIAIEKRSLQCVKLLVENGANVHARACGRFFQKGQGTCFYFGELPLSLAACTKQWDVVS 226
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               287 EDIRNLQDLTPLKLAAKEGKIEIFRHILQREFS-----------GLSHLSRKF 328
                                                                                                                                                                                                 CITY: Fort Collins
                                                                                                                                                                                                                   ADDRESSEE: Carol Talkington Verser, Ph.D. ADDRESSEE: Heska Corporation STREET: 1825 Sharp Point Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OPERATING SYSTEM: Windows 95
SOFTWARE: WordPerfect for Windows, Version 7.0
                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Verser, Carol Talkington REGISTRATION NUMBER: 37,459
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    -----NGLTPMHLCAQEDRVSVAEELVKENAAIDPKTKAGYTPLHVACHFGQINMVRFL
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                                                                                                                                   80525
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                  WordPerfect for Windows, Version 7.0
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IBM PC compatible
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US-09-065-474-33; Sequence 33; Appli; Patent No. 6063599
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INFORMATION FOR SEQ ID NO:
                             ATTORNEY/AGENT INFORMATION:
NAME: Verser, Carol Talkington
REGISTRATION NUMBER: 37,459
                                                                                                                                                                                                                             ZIP: 80525
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                        COMPUTER: IBM PC compatible OPERATING SYSTEM: Windows 95 SOTTWARE: WordPorfect for Windows, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: DIROFILARIA AND BRUGIA ANKYRIN
TITLE OF INVENTION: PROTEINS, NUCLEIC ACID MOLECULES, AND
TITLE OF INVENTION: USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Tang, Liang APPLICANT: Blehm, E. S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 37,459
REFERENCE/DOCKET NUMBER: HTTLEECOMMUNICATION INFORMATION:
TELEPHONE: 970/493-7272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
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TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      730 IE--HG-ARVSVITRASYTPLHQAAQQGHNSVVRYLLEHGASPNVHTSTGQTPLS
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                                                                                                APPLICATION NUMBER: FILING DATE: 24-APR CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                       STATE:
                                                                                                                                                                                                                                                                                                                                             CITY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: 9,0, 9505
                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                              STREET:
                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Carol Talkington
ADDRESSEE: Heska Corporation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Verser, Carol Talkington REGISTRATION NUMBER: 37,459
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FILING DATE: 24-APR-1997
                    REFERENCE/DOCKET NUMBER:
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59; Conservative
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                                                                                                                  24-APR-1998
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25.1%; Pred. No. 0.00097;
7ative 33; Mismatches 84;
                                                                                                                                                                                                                                                                                                                                                                                                    Talkington Verser,
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                                                                                                                                       US/09/065,474
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RESULT 12
US-08-623-679-9
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TELEX: 200154
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 1533 amin
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                                                                                                         REGISTRATION NUMBER: 32,983
REFERENCE/DOCKET NUMBER: 07:
TELECOMMUNICATION INFORMATION:
TELECHONE: 617,542-5070
                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/412,431
FILING DATE: 29-MAR-1995
ATTORNEY/AGENT INFORWATION:
NAME: Fasse, J. Peter
REGISTRATION NUMBER: 32,983
REGISTRATION NUMBER: 32,983
                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Boston
STATE: MA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                629 TLL---HYKANANAESKAGFTPLH--LAAQEGHREMAAL-----LIENGAKVGAQAR- 675
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                                                                                      TELEPHONE: 617/542-30
TELEPHAX: 617/542-8906
                                                                                                                                                                                                                                                                                             CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/623,679 FILING DATE: 29-MAR-1996
                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER:
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ZIP: 02110-2804
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Local Similarity 25.1%;
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TOPOLOGY: linear
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SYSTEM: PC-DOS/MS-DOS
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; TYPE: PRT; ORGANISM: Homo sapiens US-08-933-774-9
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US-08-623-679-9
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Best Local Similarity 19.9
Matches 101; Conservative
Query Match
Best Local Similarity
Matches 101; Conserv
                                                                                                                                                                                                              SOFTWARE: FastSEQ
SEQ ID NO 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: Shyjan, Andrew W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 9, Application US/08933774A Patent No. 6025137
                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE DIAGNOSIS, PREVENTION TITLE OF INVENTION: AND TREATMENT OF TUMOR PROGRESSION FILE REFERENCE: 07334/004003 CURRENT APPLICATION NUMBER: US/08/933,774A CURRENT FILING DATE: 1997-09-19 EARLIER APPLICATION NUMBER: US 08/623,679 EARLIER FILING DATE: 1996-03-29
                                                                                                                                                                                                                                                         EARLIER APPLICATION NUMBER: US 08/412,431
EARLIER FILING DATE: 1995-03-29
NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                      LENGTH: 1533
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      KIEIFRHILQREFSGLSHLSRKFTEWCYGPVRVSLYDLASVDSCEENSVLEIIAFHCKSP 365
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      --IYGEVFADQIDLYAMEINPPCGENLYDEEGKRLPPCIPGAWLTPALMACYLLVANILL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IGKMMI-DMLYFVVIMLVVLMSFGVARQAILHPEEKPSWKLARNIFYMPYWM------
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         IQKVILRDLLRFLLIYLVFLFGFAVALVSLSQEAWRPEAPTGPNATESVQPMEGQEDEGN 586
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE:
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                       3.4%;
19.9%;
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  Score 137; DB 3; Length 1533; Pred. No. 0.0011; 8; Mismatches 180; Indels 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 137; DB 1; Length 1533; Pred. No. 0.0011;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches 180;
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Conservative

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Indels 138;

Gaps

22;

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US-09-031-485-23; Sequence 23; A
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APPLICANT:
                                                                                                                                                                                                                     MEDIUM TYPE: rloppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: WordPerfect for Windows,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Tang, Liang
APPLICANT: Blehm, E. Scot
TITLE OF INVENTION: DIROFILARIA AND BRUGIA ANKYRIN
TITLE OF INVENTION: PROTEINS, NUCLEIC ACID MOLECULES,
TITLE OF INVENTION: USES THEREOF
NUMBER OF SEQUENCES: 85
CORRESPONDENCE ADDRESS:
                                                                APPLICATION NUMBER: US/08/847,429 FILING DATE: 24-APR-1997 ATTORNEY/AGENT INFORMATION:
                                                                                                                              CLASSIFICATION: 530 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      998 VNLLIAVFNNTFFEVKSISNQVWKFQR 1024
                      NAME: Verser, Carol Talkington REGISTRATION NUMBER: 37,459
                                                                                                                                                                              FILING DATE:
                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Carol Talkington Verser,
ADDRESSEE: Heska Corporation
STREET: 1825 Sharp Point Drive
CITY: Fort Collins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LNMLIALMSETVNSVATDSWSIWKLQK 664
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MIGAILRLQNQPYMGYGRVIYCVDIIFWYIRVLD---IFG-VN-----KYLGPYVMM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LFQALLTVVSQ------VLCFLAIEWYLPLLVSALVLGWLNLLYYTRGFQHTGIYSVM 526
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FGELPLSLAACTKQWD-----VVSYLLENPHQPASLQATDSQGNTVLHALVMISDNSAE
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                                                                                                                                                                                                                                                                                                                                                            80525
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  NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 28.0
Matches 73; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No. 582769
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: Tang,
APPLICANT: Blehm,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 970/484-9505
INFORMATION FOR SEQ ID NO:
TELEFAX: 9'
                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: Wordberfect for Windows,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                         TELECOMMUNICATION INFORMATION: TELEPHONE: 970/493-7272
                                                                                                                                                CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: ECORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           228
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      295 LTPLKLAAKEGKIEIFRHIL-------QREFSGLSHLSRKFTEWCYG--PVRVSLYD 342
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              189 -VHARACGRFFQKGQGTCFYF-------GELPLSLAACTKQWDVVSYLLENPHQ 234
                                                                                NAME: Verser, Carol Talkington REGISTRATION NUMBER: 37,459 REFERENCE/DOCKET NUMBER: HW-5
                                                                                                                                                                                                FILING DATE:
                                                                                                                                                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                              ZIP: 80525
                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                       STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Carol Talkington Verser,
ADDRESSEE: Heska Corporation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           77
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PLHVAA----FMGAINIVIYLLQQGANADVATVRGETPLHLAARANQTDIVRVLVRN--- 129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RGTPVDIEGKNQVTPLHVAAH 248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PASLQATDSQGNTVLHALVMISDNSAENIALVTSMYDGLLQAGARLCPTVQLEDIRNLQD
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Fort Collins
                                                                                                                                                                                                                                                                                                                                                                                                                       Colorado
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Application US/08847429A
                                                                                                                                                                                                                                                                                                                                                                                                                                                              E: Heska Corporation
1825 Sharp Point Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tang, Liang
Blehm, E. S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 303 amino acids
                  970/484-9505
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Pred. No. 0.00031; 78;
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                                                                                                                                                                                                                                                                  Version
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Вþ Qy В Qy B Q 밁

527

В

638 940

Patent No.

Qy В

792 426 738 366 702

B

Q

QΥ

Indels

76;

Gaps

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Вþ

653

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606

208

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; SEQUENCE CHARACTERISTICS:
; LENGTH: 303 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-847-429A-23
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                                                                                                                                                                                                                                                                                                                                                         Query Match 3.3%; Score 131; DB 2; Length 303; Best Local Similarity 28.0%; Pred. No. 0.00031; Matches 73; Conservative 34; Mismatches 78; Indels
                                                                                                                                                                                                                                                                                  139 LQIDRDSGNPQPLVNAQCTDDYYRGHSALHIAIEKRSLQCVKLLV------ENGAN 188
                        130 GAQVDAAARELQTPLHIASRLGN-----TDIVILLLQANA--SPNAATRDL----- 173
                                                                                                                                                                                                                                               189 -VHARACGREFQKGQGTCFYF------GELPLSLAACTKQWDVVSYLLENPHQ 234
228 RGTPVDIEGKNQVTPLHVAAH 248
                                                                                                                                                             235 PASLQATDSQGNTVLHALVMISDNSAENIALVTSMYDGLLQAGARLCPTVQLEDIRNLQD 294
                                                                                                                                                                                                               77 PLHVAA----FMGAINIVIYLLQQGANADVATVRGETPLHLAARANQTDIVRVLVRN--- 129
                                                                                                                                                                                                                                                                                                                                                          76; Gaps
                                                                                                                                                                                                                                                                                                                                                          17;
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Search completed: July 18, 2001, 15:59:03 Job time: 272 sec

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